

- (1) GENERAL INFORMATION:
- (i) APPLICANT:
 - (ii) TITLE OF INVENTION:
 - (iii) NUMBER OF SEQUENCES:
 - (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE:
 - (B) STREET:
 - (C) CITY:
 - (D) STATE:
 - (E) COUNTRY:
 - (F) ZIP:
 - (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE:
 - (B) COMPUTER:
 - (C) OPERATING SYSTEM:
 - (D) SOFTWARE:
 - (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER:
 - (B) FILING DATE:
 - (C) CLASSIFICATION:
 - (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER:
 - (B) FILING DATE:
 - (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME:
 - (B) REGISTRATION NUMBER:
 - (C) REFERENCE/DOCKET NUMBER:
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE:
 - (B) TELEFAX:
 - (C) TELEX:
 - (2) INFORMATION FOR SEQ ID NO: X:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH:
 - (B) TYPE:
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY:
 - (ii) MOLECULE TYPE:
 - (iii) ISOTOPICTICAL:
 - (iv) ANTI-SENSE:
 - (v) FRAGMENT TYPE:
 - (vi) ORIGINAL SOURCE:
 - (A) ORGANISM:
 - (B) STRAIN:
 - (C) INDIVIDUAL ISOLATE:
 - (D) DEVELOPMENTAL STAGE:
 - (E) HAPLOTYPE:
 - (F) TISSUE TYPE:
 - (G) CELL TYPE:
 - (H) CELL LINE:
 - (I) ORGANELLE:
 - (vii) IMMEDIATE SOURCE:
 - (A) LIBRARY:
 - (B) CLONE:
 - (viii) POSITION IN GENOME:
 - (A) CHROMOSOME/SEGMENT:
 - (B) MAP POSITION:
 - (C) UNITS:
 - (ix) FEATURE:
 - (A) NAME/KEY:
 - (B) LOCATION:
 - (C) IDENTIFICATION METHOD:
 - (D) OTHER INFORMATION:
 - (x) PUBLICATION INFORMATION:
 - (A) AUTHORS:
 - (B) TITLE:
 - (C) JOURNAL:
 - (D) VOLUME:
 - (E) ISSUE:
 - (F) PAGES:
 - (G) DATE:
 - (H) DOCUMENT NUMBER:
 - (I) FILING DATE:
 - (J) PUBLICATION DATE:
 - (3) RELEVANT RESIDUES:
 - (4) SEQUENCE DESCRIPTION: SEQ ID NO: X:

Mandatory information is missing.

See § 1.823(b)(1) of the old sequence rules

(2) INFORMATION FOR SEQ ID NO:1:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 27 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Oligonucleotides

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

CGTGCTACTT CCATTGTCA CGTCCTG

27

(2) INFORMATION FOR SEQ ID NO:2:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 21 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Oligonucleotides

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

TGCCGTGATA GCAAATTGA G

21

(2) INFORMATION FOR SEQ ID NO:3:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 22 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Oligonucleotides

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

AGGAAGTGAG CATTATGGAT GA

22

(2) INFORMATION FOR SEQ ID NO:4:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 20 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Oligonucleotides

**Does Not Comply
Corrected Diskette Needed**

- Please submit only one(1) file.

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

TGCTCCTCGT GCCCAGCGTT

20

(2) INFORMATION FOR SEQ ID NO:5:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 27 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Oligonucleotides

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

TTAGAATTCC CCGAGGAGGA GCTGTAC

27

(2) INFORMATION FOR SEQ ID NO:6:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 27 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Oligonucleotides

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

CTAGAATTCT GCAGTCCCAC CACCCCG

27